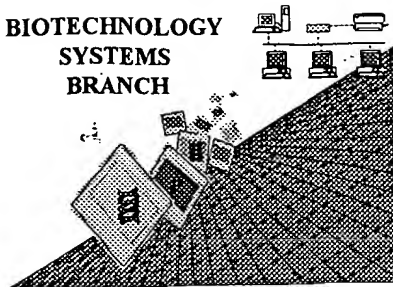


0570
1005

BIOTECHNOLOGY
SYSTEMS
BRANCH



RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/878,423
Source: OIPK
Date Processed by STIC: 8/3/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER VERSION 3.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

09/12/85 423

SEQUENCE LISTING

Does Not Comply
Corrected Diskette Needed

<110> ~~APPLICANT:~~ Aevvarsson, Arnthor
Marteinsson, Viggo T.
Hreggvidsson, Gudmundur O.
Kristjansson, Jakob K.
Fridjonsson, Olafur H.

<120> ~~TITLE:~~ Method of Obtaining Protein Diversity

<130> ~~DOCKET/FILE REFERENCE:~~ 2739.2004-001

<150> ~~PRIOR APPLICATION NUMBER:~~ Iceland 5863

<151> ~~FILING DATE:~~ 2001-02-23

<160> ~~NUMBER OF SEQUENCES:~~ 6

<170> ~~SOFTWARE:~~ FastSEQ for Windows Version 4.0

<210> ~~SEQ ID NO:~~ 1

<211> ~~LENGTH:~~ 19

<212> ~~TYPE:~~ DNA

<213> ~~ORGANISM:~~ Artificial Sequence

<220> ~~FEATURE:~~

<223> ~~OTHER INFORMATION:~~ bacterial primer

<400> ~~SEQ ID NO:~~ 1

gagtttgatc ctggctcag

19

<210> SEQ ID NO:2

<211> LENGTH: 23

<212> TYPE: DNA

<213> ORGANISM: Artificial Sequence

<220> FEATURE:

<223> OTHER INFORMATION: bacterial primer

<400> SEQ ID NO:2

gtcccagcag ccgcggtaaa tac

23

<210> SEQ ID NO:3

<211> LENGTH: 20

<212> TYPE: DNA

<213> ORGANISM: Artificial Sequence

<220> FEATURE:

<223> OTHER INFORMATION: bacterial primer

<400> SEQ ID NO:3

pp 1-2

do not include
alphabetical headings

in new sequence

Rules format. Just

use

numeric

identifiers.

edit all

sequences

09/878423 2

gactaccggg tatctaattcc

20

<210> SEQ ID NO:4

<211> LENGTH: 17

<212> TYPE: DNA

<213> ORGANISM:Artificial Sequence

<220> FEATURE:

<223> OTHER INFORMATION: bacterial primer

<400> SEQ ID NO:4

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17

<210> SEQ ID NO:5

<211> LENGTH: 38

<212> TYPE: DNA

<213> ORGANISM:Artificial Sequence

<220> FEATURE:

<223> OTHER INFORMATION: archaea primer

<400> SEQ ID NO:5

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<210> SEQ ID NO:6

<211> LENGTH: 17

<212> TYPE: DNA

<213> ORGANISM:Artificial Sequence

<220> FEATURE:

<223> OTHER INFORMATION: archaea primer

<400> SEQ ID NO:6

gacgggcggt gtgtrca

17

edit
here

sequence